REMARKS

This Amendment cancels claims 2-4 and amends claim 1, 7 and 13 in accordance with the original disclosure. In view of these amendments and the following remarks, Applicants submit that all of pending claims 1, 5-20 and 23-32 are now in condition for allowance.

Unexamined Claims

As discussed with Examiner Pak in telephone conversations on August 19, 2005 and September 15, 2005, elected claims 23-32 were not examined in the outstanding Office Action. The Examiner instructed Applicants to respond to the rejections of claims 1-20 and stated that claims 23-32 would be examined in a subsequent, non-final, Office Action. Therefore, Applicants base the above amendments and the following remarks on currently examined claims 1-20.

Claim Objections

Claims 7 and 13 stand objected to as being dependent upon a rejected base claim but would be allowable if rewritten in independent form. As set forth above, Applicants have rewritten claims 7 and 13 in independent form.

Rejections Under 35 U.S.C. § 102(b)

Claims 1-6, 8-12 and 14-20 stand rejected under 35 U.S.C. 35 U.S.C. § 102(b) for asserted anticipation by van den Beukel et al. (Applicants believe that the Examiner is referring to the van den Beukel et al. disclosure, hereinafter "van den Beukel"). The Examiner states that van den Beukel discloses chimeric receptors comprising rat alpha and Drosophila alpha subunits, chicken beta 2 subunits; and nucleic acids encoding the above receptors expressed in oocytes. Additionally, the Examiner states that the chimeric DNA were subcloned into pcDNA3 which inherently requires transformation of the host cell.

Applicants submit that the chimeric subunits disclosed by van den Beukel can be distinguished from the present invention with respect to the sequence segments which are replaced with insect sequences. Specifically, attached herewith is a sequence comparison which identifies the regions in van den Beukel's most extended replacement (designated S7.1, page 1032) in italicized type. The region defined by SEQ ID NO: 1 in Torpedo alpha 1 is written in bold and italicized. Thus, it is clear that the modified acetylcholine receptor subunit of the claimed invention comprising an α subunit of a vertebrate acetylcholine receptor in which the entire

region, which is homologous with the amino acid sequence shown in SEQ ID NO: 1, has been replaced by the corresponding region of an α subunit of an insect acetylcholine receptor is not anticipated by van den Beukel.

Applicants also respectfully submit that the Examiner is incorrect that van den Beukel discloses receptors formed from chimeric subunits containing a Beta 2 subunit. Nowhere in van den Beukel is such a description taught or suggested. Rather, van den Beukel discloses a hybrid receptor consisting of Drosophila $\alpha 2$ subunits and chick $\beta 2$ subunits (e.g., page 1032, Fig. 1). It is therefore clear that this receptor does not contain <u>chimeric</u> subunits.

Furthermore, the modified receptor subunits produced by van den Beukel are not capable of a functional response to typical acetylcholine receptor ligands such as acetylcholine or physostigmine (see page 1032, Methods and Results: "Neither ACh, Phytoestrogen nor teramethylammonium was able to induce an ion current in any of the chimeras;" also see page 1032, Conclusions: "Lack of function of the alpha7-SAD chimeras appears to be due to deficient coupling of agonist binding to gating of the ion channel..".). The chimeras disclosed by van den Beukel therefore are not suitable for a test method with which it is possible to find compounds which, as modulators, in particular as agonists or antagonists, alter the conduction properties of insect nicotinic receptors.

In contrast to van den Beukel, the chimeric subunits of the claimed invention assemble with β subunits, forming a receptor that responds with increased ion conductance to the binding of acetylcholine or imidacloprid, not only in Xenopus oocytes, but also in cell lines such as Sf9 (Examples 2 and 3 of the application).

Conclusion

In view of the above amendments and remarks, Applicants respectfully request allowance of all pending claims 1, 5-20 and 23-32.

Respectfully submitted,

Attorney for Applicants

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Attachment 2

		* 20 * 40 *	60	
a1_Human 57	:	VAKLFKDYSSVVRPVEDHRQVVEVTVGLQLIQLINVDEVNQIVTTNVRLKQGDM	VDL	:
a1_Torpedo 51	:	VANLLENYNKVIRPVEHHTHFVDITVGLQLIQLISVDEVNQIVETNVRLRQ		:
a4_Chick 51	:	LKKLFSGYNKWSRPVANISDVVLVRFGLSIAQLIDVDEKNQMMTTNVWVKQ		:
a2_Drosophila 51	:	YDDLLSNYNRLIRPVSNNTDTVLVKLGLRLSQLIDLNLKDQILTTNVWLEH		:
a2_Heliothis 51	:	YDDLLSNYNRLIRPVDKNNNTVLVKLGLRLSQLIDLNLKDQILTTNVWLEH		:
a3_Heliothis	:	YDDLLSNYNRLIRPVTNVSDILTVRLGLKLSQLMEVNLKNQVMTTNLWVEQ		:
a2_Myzus 51	:	YDDLLSNYNRLIRPVGNNSDRLTVKMGLKLSQIIEVNLRNQIMTTNVWVEQ		:
a3_Drosophila	:	YDDLLSNYNKLVRPVVNVTDALTVRIKLKLSQLIDVNLKNQIMTTNLWVEQ		:
a1_Heliothis	:	YDDLLSNYNKLVRPVLNVSDALTVRIKLKLSQLIDVNLKNQIMTTNLWVEQ		:
a3_Myzus 51	:	YDDLLSNYNKLVRPVLNNTDPLPVRIKLKLSQLIDINLKNQIMTTNLWVEQ		:
a7_Rat	:	RRLYKELVKNYNPLERPVANDSQPLTVYFSLSLLQIMDVDEKNQVLTTNIWLQM		:
54				
54				
54		* 80 * 100 *	120	
54 a1_Human 117	:	* 80 * 100 * PRPSCVTLGVPLFSHLQNEQWVDYNLKWNPDDYGGVKKIHIPSEKIWRPDLVLYNNA	120 DGD	:
al_Human	:		DGD	:
al_Human 117 al_Torpedo	:	PRPSCVTLGVPLFSHLQNEQWVDYNLKWNPDDYGGVKKIHIPSEKIWRPDLVLYNNA	DGD	:
a1_Human 117 a1_Torpedo 92 a4_Chick	: : :	PRPSCVTLGVPLFSHLQNEQWVDYNLKWNPDDYGGVKKIHIPSEKIWRPDLVLYNNA	DGD DGD DGD	: : :
a1_Human 117 a1_Torpedo 92 a4_Chick 92 a2_Drosophila	: : : : :	PRPSCVTLGVPLFSHLQNEQWVDYNLKWNPDDYGGVKKIHIPSEKIWRPDLVLYNNAQWIDVRLRWNPADYGGIKKIRLPSDDVWLPDLVLYNNAEWHDYKLRWDPQEYENVTSIRIPSELIWRPDIVLYNNA	DGD DGD DGD	: : :
a1_Human 117 a1_Torpedo 92 a4_Chick 92 a2_Drosophila 92 a2_Heliothis 92 a3_Heliothis	: : : : :	PRPSCVTLGVPLFSHLQNEQWVDYNLKWNPDDYGGVKKIHIPSEKIWRPDLVLYNNAQWIDVRLRWNPADYGGIKKIRLPSDDVWLPDLVLYNNAEWHDYKLRWDPQEYENVTSIRIPSELIWRPDIVLYNNAEWQDHKFKWDPSEYGGVTELYVPSEHIWLPDIVLYNNA	DGD DGD DGD DGE DGE	:
a1_Human 117 a1_Torpedo 92 a4_Chick 92 a2_Drosophila 92 a2_Heliothis 92 a3_Heliothis 92 a2_Myzus	: : : : : :	PRPSCVTLGVPLFSHLQNEQWVDYNLKWNPDDYGGVKKIHIPSEKIWRPDLVLYNNAQWIDVRLRWNPADYGGIKKIRLPSDDVWLPDLVLYNNAEWHDYKLRWDPQEYENVTSIRIPSELIWRPDIVLYNNAEWQDHKFKWDPSEYGGVTELYVPSEHIWLPDIVLYNNAEWEDHKFKWDPLEYGGVKELYVPSEHIWLPDIVLYNNA	DGD DGD DGD DGE DGE DGE	:
a1_Human 117 a1_Torpedo 92 a4_Chick 92 a2_Drosophila 92 a2_Heliothis 92 a3_Heliothis 92 a2_Myzus 92 a3_Drosophila	: : : : : : : : : : : : : : : : : : : :	PRPSCVTLGVPLFSHLQNEQWVDYNLKWNPDDYGGVKKIHIPSEKIWRPDLVLYNNAQWIDVRLRWNPADYGGIKKIRLPSDDVWLPDLVLYNNAEWHDYKLRWDPQEYENVTSIRIPSELIWRPDIVLYNNAEWQDHKFKWDPSEYGGVTELYVPSEHIWLPDIVLYNNAEWEDHKFKWDPLEYGGVKELYVPSEHIWLPDIVLYNNAKWFDYKLQWNPDDYGGVEMLYVPSEHIWLPDIVLYNNW	DGD DGD DGD DGE DGE DGE	:
al_Human 117 al_Torpedo 92 a4_Chick 92 a2_Drosophila 92 a2_Heliothis 92 a3_Heliothis 92 a2_Myzus 92 a3_Drosophila 92 a1_Heliothis	: : : : : : : : : : : : : : : : : : : :	PRPSCVTLGVPLFSHLQNEQWVDYNLKWNPDDYGGVKKIHIPSEKIWRPDLVLYNNAQWIDVRLRWNPADYGGIKKIRLPSDDVWLPDLVLYNNAEWHDYKLRWDPQEYENVTSIRIPSELIWRPDIVLYNNAEWQDHKFKWDPSEYGGVTELYVPSEHIWLPDIVLYNNAEWEDHKFKWDPLEYGGVKELYVPSEHIWLPDIVLYNNAKWFDYKLQWNPDDYGGVEMLYVPSEHIWLPDIVLYNNW	DGD DGD DGD DGE DGE DGN DGN DGN	:
a1_Human 117 a1_Torpedo 92 a4_Chick 92 a2_Drosophila 92 a2_Heliothis 92 a3_Heliothis 92 a2_Myzus 92 a3_Drosophila 92	: : : : : : : :	PRPSCVTLGVPLFSHLQNEQWVDYNLKWNPDDYGGVKKIHIPSEKIWRPDLVLYNNAQWIDVRLRWNPADYGGIKKIRLPSDDVWLPDLVLYNNAEWHDYKLRWDPQEYENVTSIRIPSELIWRPDIVLYNNAEWQDHKFKWDPSEYGGVTELYVPSEHIWLPDIVLYNNAEWEDHKFKWDPLEYGGVKELYVPSEHIWLPDIVLYNNAKWFDYKLQWNPDDYGGVEMLYVPSEHIWLPDIVLYNNWEWNDYKLKWNPEDYGGVDTLHVPSEHIWLPDIVLYNNASWYDYKLKWEPKEYGGVEMLHVPSDHIWRPDIVLYNNA	DGD DGD DGE DGE DGN DGN DGN DGN	: : : : :



				-					
			*	140	* .	160	*	180	
al_Human 177	:	FAIVKFTKV	/LLQYT0	SHITWTPPAI	FKSYCEIIVI	THFPFDEQNC	SMKLGTWTYI	GSVVAIN	:
a1_Torpedo 152	:	FAIVHMTK1	LLLDYTC	KIMWTPPAI	FKSYCEIIV	THEPEDQQNC	<u>T</u> MKLGIWTYI	GTKVSIS	:
a4_Chick 152	:	FAVTHLTKA	HLFYDO	RIKWMPPAI	YKSSCSIDVI	TFFPFDQQNC	KMKFGSWTYI	KAKIDLV	:
a2_Drosophila 152	:	YVVTTMTKA	AILHYTG	KVVWTPPAI	FKSSCEIDVF	RYFPFDQQTC	FMKFGSWTYE	GDQIDLK	:
a2_Heliothis 152	:	YVVTTMTKA	AVLHHTG	KVLWTPPAI	FKSSCEIDVE	RYFPFDQQTC	FLKFGSWSYD	GDQIDLK	:
a3_Heliothis	:	YEVTLMTKA	ATLKYTO	EVNWKPPAI	YKSSCEINVE	EYFPFDEQTC	FMKFGSWTYN	IGAQVDLK	:
a2_Myzus 152	:	YEVTIMTKA	AILHYTG	KVVWKPPAI	YKSFCEINVE	EYFPFDEQTC	SMKFGSWTYD	GYMMDLR	:
a3_Drosophila	:	FEVTLATKA	ATLNYTO	RVEWRPPAI	YKSSCEIDVE	EYFPFDEQTC	VMKFGSWTYD	GFQVDLR	:
al_Heliothis	:	FEVTLATKA	ATLNYTO	RVEWRPPAI	YKSSCEIDVE	EYFPFDQQTC	VMKFGSWTYD	GFQVDLR	:
a3_Myzus	:	FEVTLATKA	MLHYSC	RVEWKPPAI	YKSSCEIDVE	EFFPFDEQTC	VMKFGSWTYD	GFQVDLR	:
a7_Rat 155	:	FDATFHTNV	/LVNASG	HCQYLPPGI	FKSSCYIDVF	RWFPFDVQQC	KLKFGSWSYC	GWSLDLQ	:
	-			200		222		0.10	
al_Human		PESDQP	*	200 -DLSNFMES	* GEWVIKESRG	220 GWKHSVTYSC	* CPDTPYLDII	240 YHFVMQR	: ,
al_Human 227 al_Torpedo		PESDQP	* 	-DLSNFMES		SWKHSVTYSC	* CPDTPYLDII CPDTPYLDII	YHFVMQR	: .
al_Human 227 al_Torpedo 202 a4_Chick	:		*	-DLSNFMES	GEWVMKDYRG	SWKHSVTYSC		YHFVMQR	-
al_Human 227 al_Torpedo 202 a4_Chick 201 a2_Drosophila	:	PESDRP		-DLSNFMES -DLSTFMES -DQLDYWES	GEWVMKDYRG GEWVIINAVG	GWKHSVTYSC GWKHWVYYTC GNYNSKKYEC	CPDTPYLDIT	YHFVMQR YHFIMQR YSFIIRR	:
al_Human 227 al_Torpedo 202 a4_Chick 201 a2_Drosophila 211 a2_Heliothis	:	PESDRP SMHSHV HISQKNDKD	ONKVEIG	-DLSNFMES -DLSTFMES -DQLDYWES IDLREYYPS	GEWVMKDYRG GEWVIINAVG VEWDILGVPA	SWKHSVTYSC SWKHWVYYTC SNYNSKKYEC AERHEKYYPC	CPDTPYLDIT	YHFVMQR YHFIMQR YSFIIRR FNITLRR	:
al_Human 227 al_Torpedo 202 a4_Chick 201 a2_Drosophila 211 a2_Heliothis 209 a3_Heliothis	:	PESDRP SMHSHV HISQKNDKD	DMVDVG	-DLSNFMES -DLSTFMES -DQLDYWES IDLREYYPS	GEWVMKDYRG GEWVIINAVG VEWDILGVPA	SWKHSVTYSC SWKHWVYYTC SNYNSKKYEC AERHEKYYPC AERHERYYPC	CPDTPYLDIT CTEI-YPDIT CAEP-YPDIF	YHFVMQR YHFIMQR YSFIIRR FNITLRR FNITLRR	:
al_Human 227 al_Torpedo 202 a4_Chick 201 a2_Drosophila 211 a2_Heliothis 209 a3_Heliothis 210 a2_Myzus	: : : : : : : : : : : : : : : : : : : :	PESDRP SMHSHV HISQKNDKD HINQKKG	ONKVEIG GDMVDVG	-DLSNFMES -DLSTFMES -DQLDYWES IDLREYYPS IDLREYYPS	GEWVMKDYRG GEWVIINAVG VEWDILGVPA VEWDILGVPA	SWKHSVTYSC SWKHWVYYTC SNYNSKKYEC AERHEKYYPC AERHERYYPC ATRNEEYYPC	CPDTPYLDIT CTEI-YPDIT CAEP-YPDIF CQEP-YPDIF	YHFVMQR YHFIMQR YSFIIRR FNITLRR FNITLRR FKLTMRR	:
al_Human 227 al_Torpedo 202 a4_Chick 201 a2_Drosophila 211 a2_Heliothis 209 a3_Heliothis 210 a2_Myzus 210 a3_Drosophila	: : : : : : : : : : : : : : : : : : : :	PESDRP SMHSHV HISQKNDKD HINQKKG HMDQSP-GS HISQAP-DS	DNKVEIG SDMVDVG SSLVHVG	-DLSNFMES -DLSTFMES -DQLDYWES IDLREYYPS IDLREYYPS IDLSEFYLS	GEWVMKDYRG GEWVIINAVG VEWDILGVPA VEWDILGVPA VEWDILEVPA VEWDIMGVPA	SWKHSVTYSC SWKHWVYYTC SNYNSKKYEC AERHEKYYPC AERHERYYPC ATRNEEYYPC	CPDTPYLDIT CTEI-YPDIT CAEP-YPDIF CQEP-YPDIF CPEP-FSDIT	YHFVMQR YHFIMQR YSFIIRR FNITLRR FNITLRR FKLTMRR FKLTMRR	:
al_Human 227 al_Torpedo 202 a4_Chick 201 a2_Drosophila 211 a2_Heliothis 209 a3_Heliothis 210 a2_Myzus 210 a3_Drosophila 210 a1_Heliothis	: : : : : : : : : : : : : : : : : : : :	PESDRP SMHSHV HISQKNDKD HINQKKG HMDQSP-GS HISQAP-DS HIDELN-GT	ONKVEIG SDMVDVG SSLVHVG	-DLSNFMES -DLSTFMES -DQLDYWES IDLREYYPS IDLSEFYLS IDLSEFYLS IDLQDYYLS	GEWVMKDYRG GEWVIINAVG VEWDILGVPA VEWDILEVPA VEWDIMGVPA	SWKHSVTYSC SWKHWVYYTC SNYNSKKYEC AERHEKYYPC AERHERYYPC ATRNEEYYPC AVRHEKFYVC	CPDTPYLDIT CTEI-YPDIT CAEP-YPDIF CQEP-YPDIF CPEP-FSDIT CEEP-YLDIF	YHFVMQR YHFIMQR YSFIIRR FNITLRR FNITLRR FKLTMRR FNITLRR	: : : : : : : : : : : : : : : : : : : :
al_Human 227 al_Torpedo 202 a4_Chick 201 a2_Drosophila 211 a2_Heliothis 209 a3_Heliothis 210 a2_Myzus 210 a3_Drosophila 210		PESDRP SMHSHV HISQKNDKD HINQKKG HMDQSP-GS HISQAP-DS HIDELN-GT	DNKVEIG SDMVDVG SDVIEVG TNVVEVG	-DLSNFMES -DLSTFMES -DQLDYWES IDLREYYPS IDLSEFYLS IDLODYYLS VDLSEFYTS	GEWVMKDYRG GEWVIINAVG VEWDILGVPA VEWDILEVPA VEWDIMGVPA VEWDILEVPA	SWKHSVTYSC SWKHWVYYTC SNYNSKKYEC AERHEKYYPC AERHERYYPC ATRNEEYYPC AVRHEKFYVC AVRNEKFYTC	CPDTPYLDIT CTEI-YPDIT CAEP-YPDIF CQEP-YPDIF CPEP-FSDIT CEEP-YLDIF	YHFVMQR YHFIMQR YSFIIRR FNITLRR FNITLRR FKLTMRR FNITLRR FNITLRR	: : : : :